

GenCode version 4.5  
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### 1M protein - protein search, using sw model

Run (n): March 14, 2002, 17:15:02 - March 14, 2002, 17:43:56 (0.47 hours)

$$115 = (14 - 2) \cdot 8 + (10 \cdot 6 - 7)$$

Sequencia: 1 MAMC/2004/00550 14

printing table: HLC51062

**Searched:** 100056 soaps, 36664827 residues

Total number of hits satisfying chosen parameters. 100055

Minimum LB seq length: 1

Post processing: Minimum Match

Listed first 45 summaries

Out class : **SwissProt\_39:\***

Evaluation :

pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

[illegible][illegible][illegible]





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EMBL: Molecular Biology

Accession: P000129; P001001; P001002

Accession: P000129; P001001; P001002

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Query Match: 51.1% Score: 381; 108 12; Length: 514  
Best Local Similarity: 60.0% (Prod. No. 47)  
Matches: 62 Conserved 20 Mismatches: 17 Indels: 0 Gaps: 0

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 001 Spermatozoa: Mammalia: Eutheria: Carnivora: Canidae: Felidae: Felis tigris.  
 002 Asteraceae: Asteraceae: Helianthus: Helianthus: Helianthus.  
 003 NCBI\_TaxID: 4043.  
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 001 Spermatozoa: Mammalia: Eutheria: Carnivora: Canidae: Felidae: Felis tigris.  
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